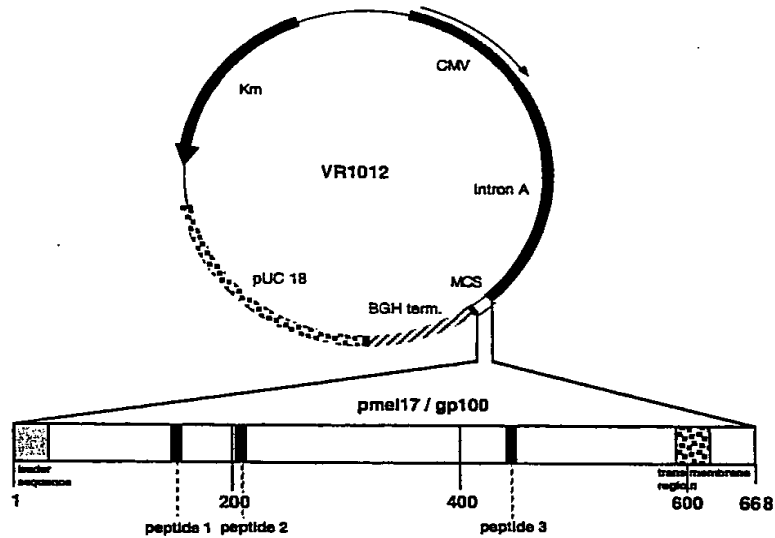


A



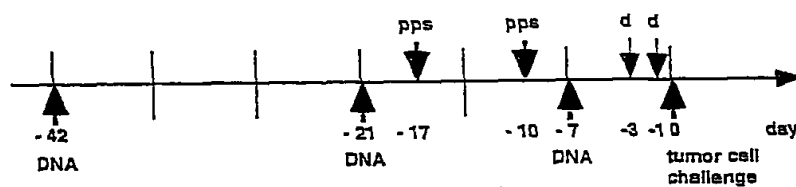
B

human	1	MDLVLRCLLEPAVIGALLAVGATKVPNRQDWLGVSRLRTRAWNRQLYPEWTEAQRLLC
mouse	1	MVGQRRSEFLPVILVLSALLAVGALECSRNDQWLGVPRLVTKTNRQLYPEWTEVQSGNC
human	61	WRGGQVSLKVSNDGPTLLGANASFSTALNPPGSKQVLPDGGQVWVNNNTIINGSQVWGGQF
mouse	61	WRGGQVSLRVNDGPTLLGANASFSTALNPPGSKQVLPDGGQVWVNNNTIINGSQVWGGQF
human	121	VYPQETDDACTEFDGGPCPSCSWSCRRSEFVYVWKTWCQYQVLLGGFVSCISIGTCRAMLC
mouse	121	VYPQETDDACVFEFDGGPCPSCFKPKRRSEFVYVWKTWCQYQVLLGGFVSRSSIATRHAKLG
human	181	THTMEVTVYHRRGSRSYVPLAHSSAFTITDQVPFVSVSQDRAIDCGNKHFLRNQPLTF
mouse	181	THTMEVTVYHRRGSCSYVPLAHASSFTITDQVPFVSVSQDRAIDGCKHFLRNQPLTF
human	241	ALQLHDPSCYLAEADLSYTWDFGDSGGLISRAFPVTHTYLESQSVTAQVVLQAAIPILAS
mouse	241	ALQLHDPSCYLAEADLSYTWDFGDCGGLISRAFDVTHTYLESQSVTAQVVLQAAIPILVS
human	301	CGSSRPVPGTIDGHRPTAEAPNTIAGOVPTDEWCTTPCOAPTAPSGTTSVQVPTTEVLS
mouse	301	CGSSRPVPGTIDGHRPTAEAPNTISROCTTKRVVGTTPCOMPTOPSGTTVQVPTTEVTA
human	361	TAFTVQNPTESTGCTEKVPVSEVMCTTLAEMSTPEATGCTEALVSIVVLSGTTAAQVTE
mouse	361	TTSEOMLT-----SAVVDITLAEVSTPESTGCTETWRP-----SGTTVAQATE
human	421	TEWVETTARELPPIPEPEGPDASSITSTESTIGSTCPILLDCTATIRLVKQVPLDCVLYRY
mouse	403	TE-----GPDASPIPTOSTIGSISPLLEDCTATIRLVKQVPLDCVLYRY
human	481	GSFSVLDIVQGISSAEILQAVFSEGDFAFELTVSCQGGLPKEACMEISSPGCOPPAQRL
mouse	448	GSFSVLDIVQGISSAEILQAVFSEGDFAFELTVSCQGGLPKEACMDISSPGCOPPAQRL
human	541	QCPVLPSPACQLVLHQILKGGSCITYCLNVSLADANSLAVVSTQLIMPVPGTILLTQQAAGL
mouse	508	QCSVPSPSPCQLVLHQVLKGGSCITYCLNVSLADANSLAVASTQLVVE-----GQDGGI
human	601	GQVPLIVGILLVILMAVVLASLIY-----RRRLMKQDFSVQDPHSSSEWLRTPRIFCSCEI
mouse	561	GCAPLEVIGILLVILMAVVLASLIY-----GIDLRSAQFPKCEVALTAAPASGLRA-R-----G-L
human	657	GENSPILLSGQQV
mouse	615	GENSPILLSGQQV

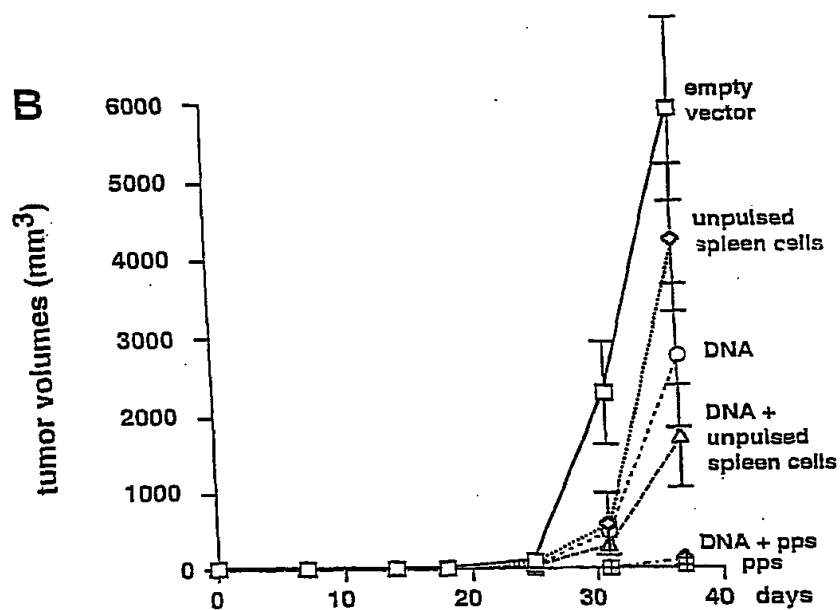
Fig. 1

A prophylactic

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B



C

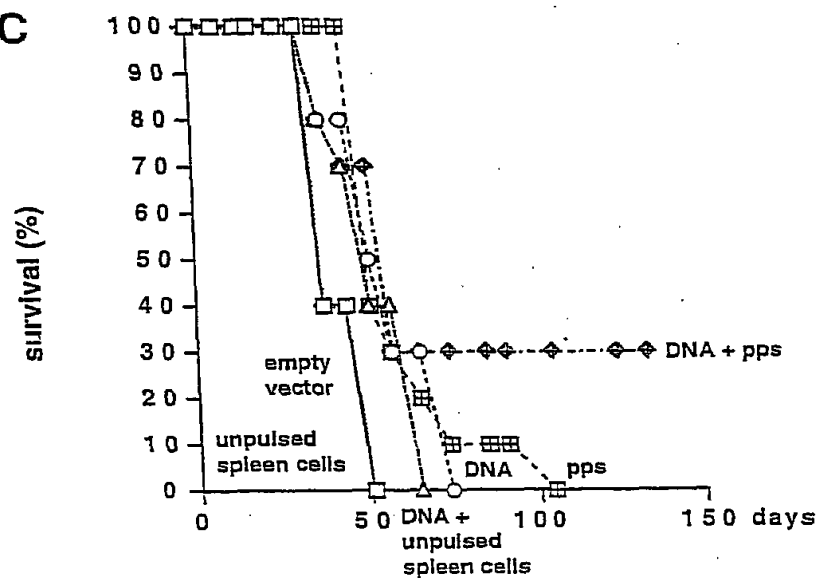


Fig. 2

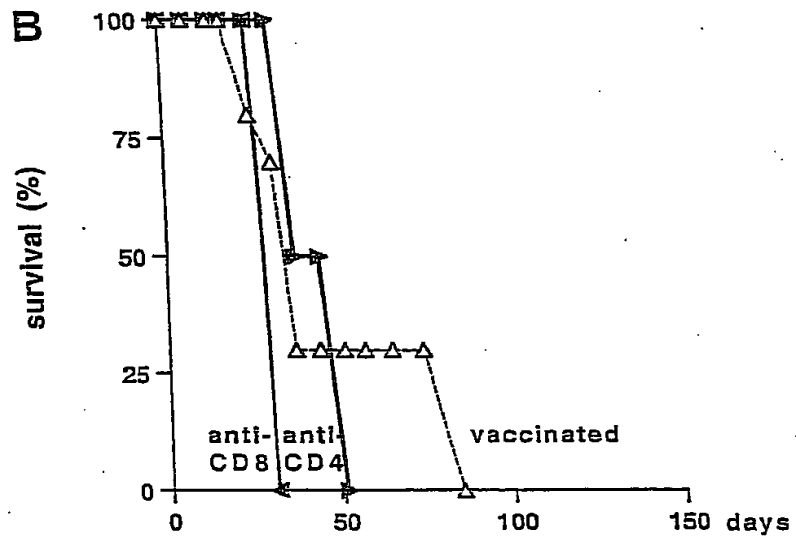
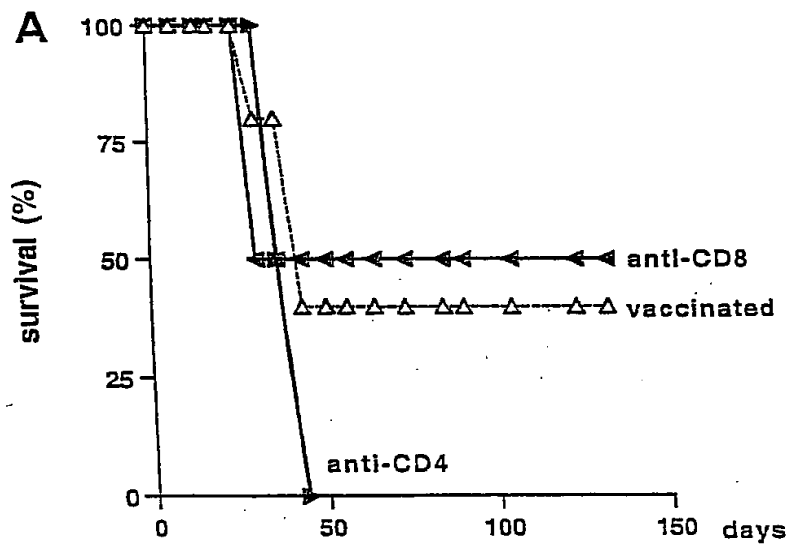


Fig. 3

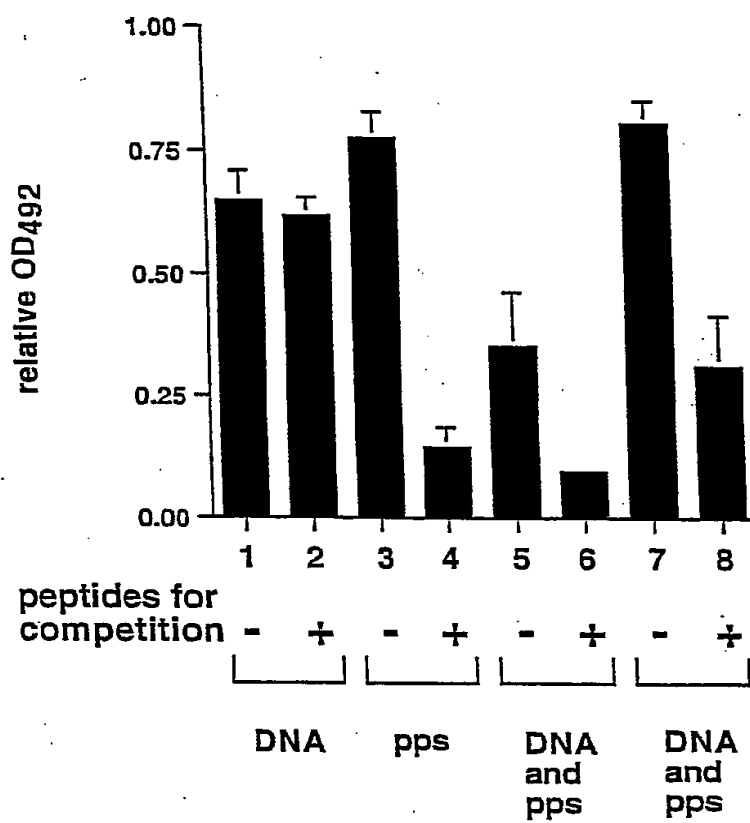


Fig. 4

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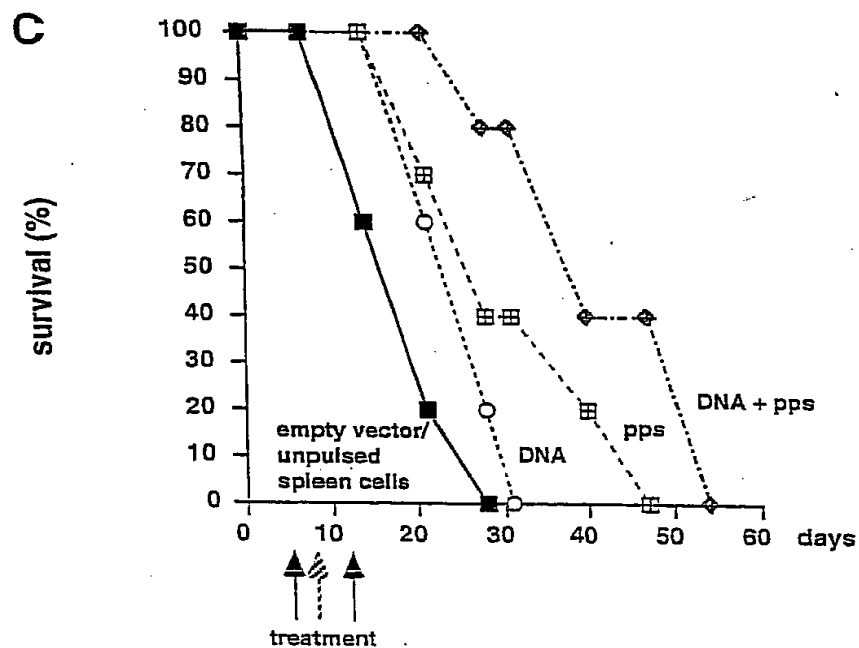
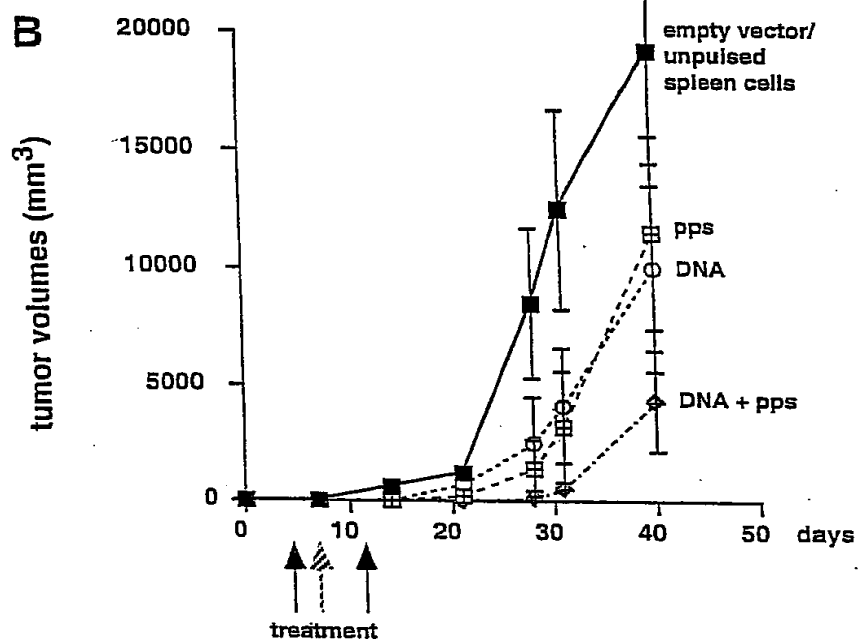
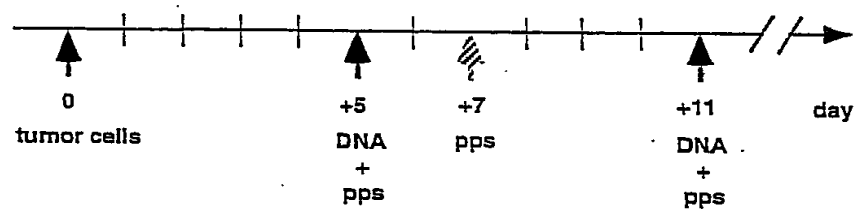


Fig. 5